

# Unsupervised Learning in Python

## 2). Visualization with hierarchical clustering and t-SNE:

### a). Hierarchical Clustering of the grain data

# Perform the necessary imports

```
from scipy.cluster.hierarchy import dendrogram, linkage
```

```
import matplotlib.pyplot as plt
```

# Calculate the linkage: mergings

```
mergings = linkage(samples, method='complete')
```

# Plot the dendrogram, using varieties as labels

```
dendrogram(mergings,
```

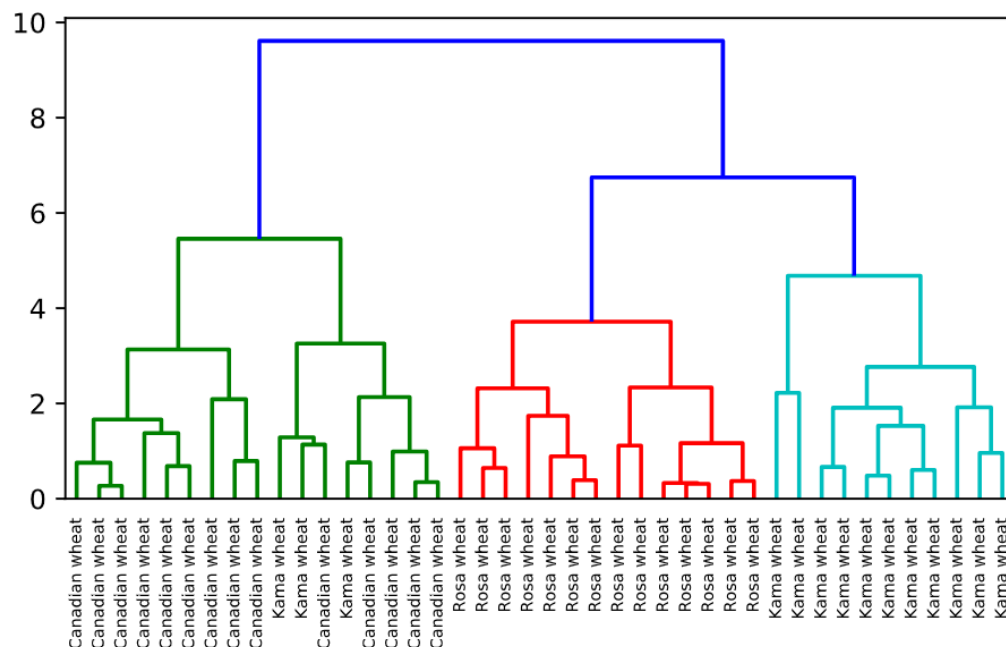
```
    labels=varieties,
```

```
    leaf_rotation=90,
```

```
    leaf_font_size=6,
```

```
)
```

```
plt.show()
```



b). Hierarchies of Stocks

# Import normalize

from sklearn.preprocessing import normalize

# Normalize the movements: normalized\_movements

normalized\_movements = normalize(movements)

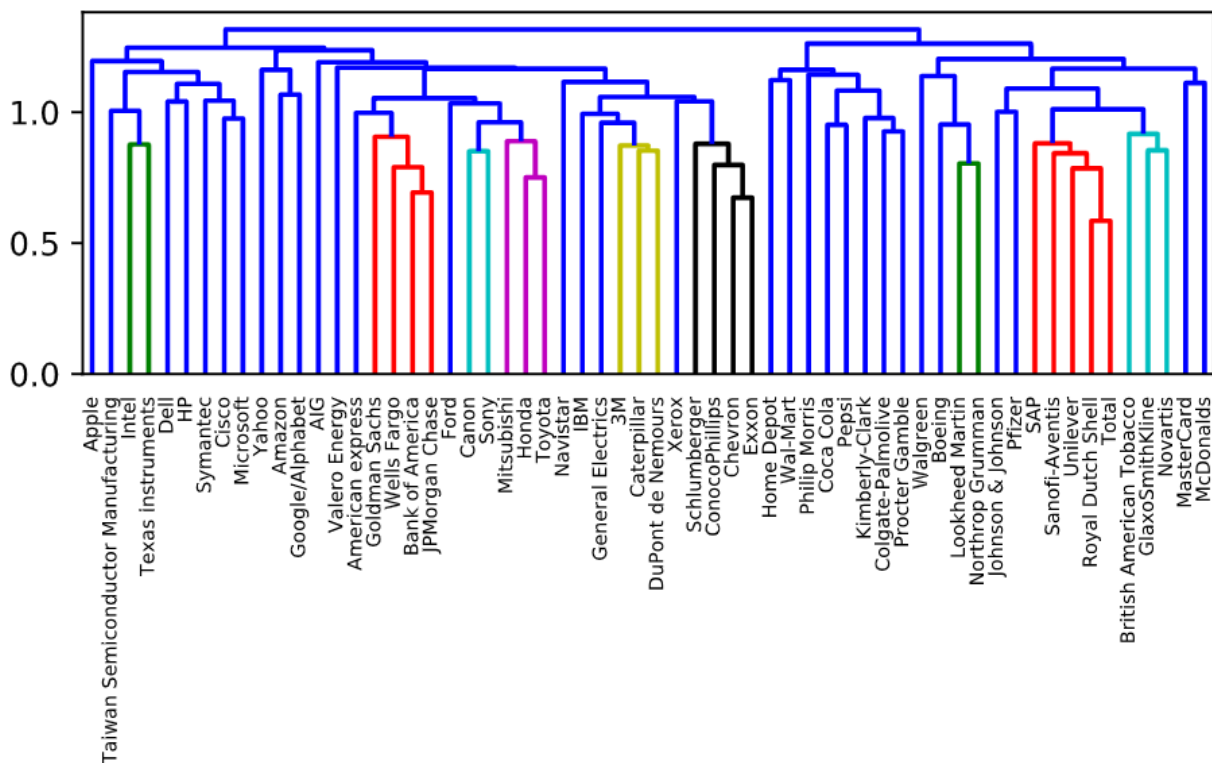
# Calculate the linkage: mergings

mergings = linkage(normalized\_movements,'complete')

# Plot the dendrogram

dendrogram(mergings, labels=companies, leaf\_rotation=90, leaf\_font\_size=6)

plt.show()



c). Different linkage, different hierarchical clustering!

# Perform the necessary imports

import matplotlib.pyplot as plt

from scipy.cluster.hierarchy import linkage, dendrogram

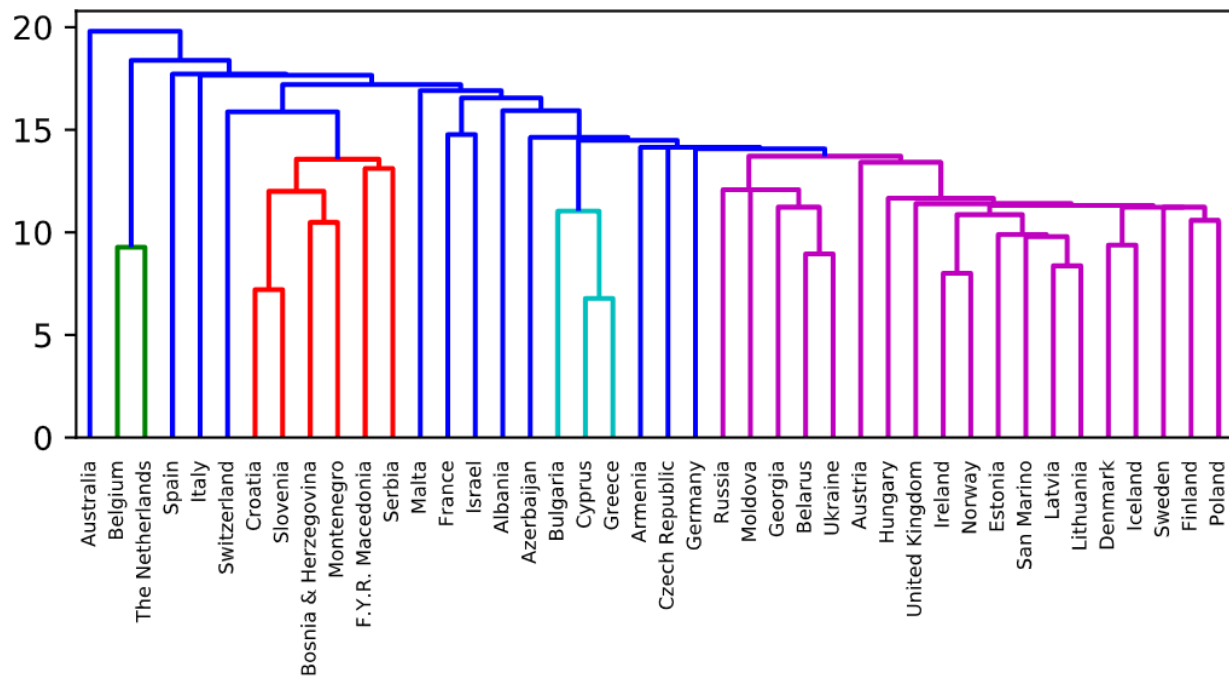
# Calculate the linkage: mergings

mergings = linkage(samples, method='single')

# Plot the dendrogram

dendrogram(mergings, labels=country\_names, leaf\_rotation=90, leaf\_font\_size=6)

plt.show()



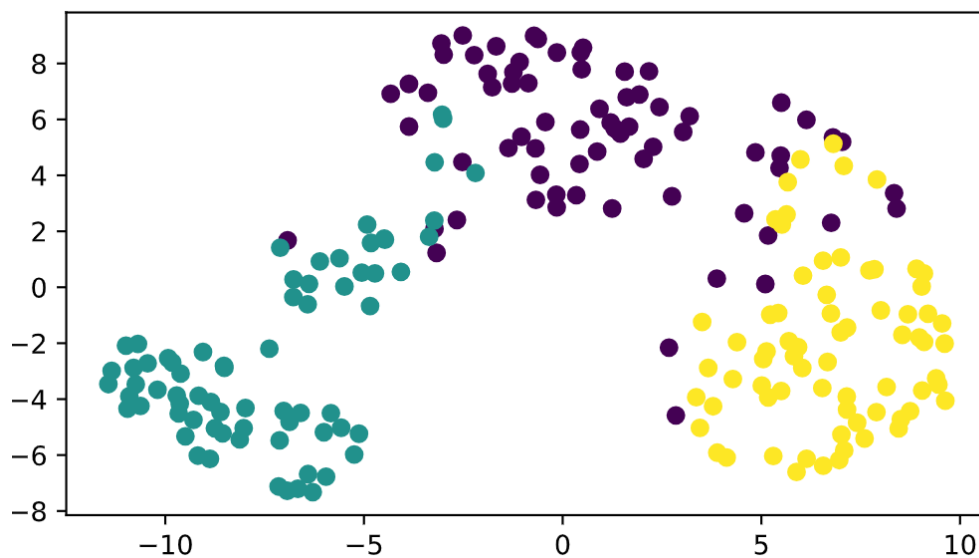
**d). Extracting the cluster labels****# Perform the necessary imports****import pandas as pd****from scipy.cluster.hierarchy import fcluster****# Use fcluster to extract labels: labels****labels = fcluster(mergings,6 , criterion='distance')****# Create a DataFrame with labels and varieties as columns: df****df = pd.DataFrame({'labels': labels, 'varieties': varieties})****# Create crosstab: ct****ct = pd.crosstab(df['labels'], df['varieties'])****# Display ct****print(ct)**

&lt;script.py&gt; output:

varieties Canadian wheat Kama wheat Rosa wheat

labels

1	14	3	0
2	0	0	14
3	0	11	0

**e). t-SNE Visualization of grain dataset****# Import TSNE****from sklearn.manifold import TSNE****# Create a TSNE instance: model****model = TSNE(learning\_rate=200)****# Apply fit\_transform to samples: tsne\_features****tsne\_features = model.fit\_transform(samples)****# Select the 0th feature: xs****xs = tsne\_features[:,0]****# Select the 1st feature: ys****ys = tsne\_features[:,1]****# Scatter plot, coloring by variety\_numbers****plt.scatter(xs, ys, c=variety\_numbers)****plt.show()**

```
# Import TSNE
```

## # Create a TSNE instance: model

```
# Apply fit_transform to normalized_movements: tsne_features
```

### # Select the 0th feature: xs

```
# Select the 1th feature: ys
```

```
# Scatter plot
```

### # Annotate the points

```
plt.annotate(company, (x, y), fontsize=5, alpha=0.75)
```

**plt.show()**